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## SEQUENCE LISTING

<110> PAKOLA, STEVE  
DE SMET, MARK

<120> PHARMACOLOGICAL VITREOLYSIS

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<140> 10/729,475

<141> 2003-12-05

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<151> 2002-12-06

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<170> PatentIn Ver. 3.2

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15

48

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35 40 45	
acc ttg ata tcc cca gag tgg gtg ttg act gct gcc cac tgc ttg gag Thr Leu Ile Ser Pro Glu Trp Val Leu Thr Ala Ala His Cys Leu Glu	192
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65 70 75 80	
gaa gtg aat ctc gaa ccg cat gtt cag gaa ata gaa gtg tct agg ctg Glu Val Asn Leu Glu Pro His Val Gln Glu Ile Glu Val Ser Arg Leu	288
85 90 95	
ttc ttg gag ccc aca cga aaa gat att gcc ttg cta aag cta agc agt Phe Leu Glu Pro Thr Arg Lys Asp Ile Ala Leu Leu Lys Leu Ser Ser	336
100 105 110	
cct gcc gtc atc act gac aaa gta atc cca gct tgt ctg cca tcc cca Pro Ala Val Ile Thr Asp Lys Val Ile Pro Ala Cys Leu Pro Ser Pro	384
115 120 125	
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130 135 140	
gaa acc caa ggt act ttt gga gct ggc ctt ctc aag gaa gcc cag ctc Glu Thr Gln Gly Thr Phe Gly Ala Gly Leu Leu Lys Glu Ala Gln Leu	480
145 150 155 160	
cct gtg att gag aat aaa gtg tgc aat cgc tat gag ttt ctg aat gga Pro Val Ile Glu Asn Lys Val Cys Asn Arg Tyr Glu Phe Leu Asn Gly	528
165 170 175	
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180 185 190	
gac agt tgc cag ggt gac agt gga ggt cct ctg gtt tgc ttc gag aag Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Phe Glu Lys	624
195 200 205	
gac aaa tac att tta caa gga gtc act tct tgg ggt ctt ggc tgt gca Asp Lys Tyr Ile Leu Gln Gly Val Thr Ser Trp Gly Leu Gly Cys Ala	672
210 215 220	
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 35 40 45

Thr Leu Ile Ser Pro Glu Trp Val Leu Thr Ala Ala His Cys Leu Glu  
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Lys Ser Pro Arg Pro Ser Ser Tyr Lys Val Ile Leu Gly Ala His Gln  
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Glu Val Asn Leu Glu Pro His Val Gln Glu Ile Glu Val Ser Arg Leu  
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Phe Leu Glu Pro Thr Arg Lys Asp Ile Ala Leu Leu Lys Leu Ser Ser  
 100 105 110

Pro Ala Val Ile Thr Asp Lys Val Ile Pro Ala Cys Leu Pro Ser Pro  
 115 120 125

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Pro Val Ile Glu Asn Lys Val Cys Asn Arg Tyr Glu Phe Leu Asn Gly  
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Arg Val Gln Ser Thr Glu Leu Cys Ala Gly His Leu Ala Gly Gly Thr  
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Asp Lys Tyr Ile Leu Gln Gly Val Thr Ser Trp Gly Leu Gly Cys Ala  
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Glu Asp Cys Met Phe Gly Asn Gly Lys Gly Tyr Arg Gly Lys Arg Ala
      20           25           30

acc act gtt act ggg acg cca tgc cag gac tgg gct gcc cag gag ccc 144
Thr Thr Val Thr Gly Thr Pro Cys Gln Asp Trp Ala Ala Gln Glu Pro
      35           40           45

cat aga cac agc att ttc act cca gag aca aat cca cgg gcg ggt ctg 192
His Arg His Ser Ile Phe Thr Pro Glu Thr Asn Pro Arg Ala Gly Leu
      50           55           60

gaa aaa aat tac tgc cgt aac cct gat ggt gat gta ggt ggt ccc tgg 240
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Gln Cys Ala Ala Pro Ser Phe Asp Cys Gly Lys Pro Gln Val Glu Pro			
100	105	110	
aag aaa tgt cct gga agg gtt gtg ggg ggg tgt gtg gcc cac cca cat		384	
Lys Lys Cys Pro Gly Arg Val Val Gly Gly Cys Val Ala His Pro His			
115	120	125	
tcc tgg ccc tgg caa gtc agt ctt aga aca agg ttt gga atg cac ttc		432	
Ser Trp Pro Trp Gln Val Ser Leu Arg Thr Arg Phe Gly Met His Phe			
130	135	140	
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Cys Gly Gly Thr Leu Ile Ser Pro Glu Trp Val Leu Thr Ala Ala His			
145	150	155	160
tgc ttg gag aag tcc cca agg cct tca tcc tac aag gtc atc ctg ggt		528	
Cys Leu Glu Lys Ser Pro Arg Pro Ser Ser Tyr Lys Val Ile Leu Gly			
165	170	175	
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Ala His Gln Glu Val Asn Leu Glu Pro His Val Gln Glu Ile Glu Val			
180	185	190	
tct agg ctg ttc ttg gag ccc aca cga aaa gat att gcc ttg cta aag		624	
Ser Arg Leu Phe Leu Glu Pro Thr Arg Lys Asp Ile Ala Leu Leu Lys			
195	200	205	
cta agc agt cct gcc gtc atc act gac aaa gta atc cca gct tgt ctg		672	
Leu Ser Ser Pro Ala Val Ile Thr Asp Lys Val Ile Pro Ala Cys Leu			
210	215	220	
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Pro Ser Pro Asn Tyr Val Val Ala Asp Arg Thr Glu Cys Phe Ile Thr			
225	230	235	240
ggc tgg gga gaa acc caa ggt act ttt gga gct ggc ctt ctc aag gaa		768	
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245	250	255	
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His Arg His Ser Ile Phe Thr Pro Glu Thr Asn Pro Arg Ala Gly Leu  
50 55 60

Glu Lys Asn Tyr Cys Arg Asn Pro Asp Gly Asp Val Gly Gly Pro Trp  
65 70 75 80

Cys Tyr Thr Thr Asn Pro Arg Lys Leu Tyr Asp Tyr Cys Asp Val Pro  
85 90 95

Gln Cys Ala Ala Pro Ser Phe Asp Cys Gly Lys Pro Gln Val Glu Pro  
100 105 110

Lys Lys Cys Pro Gly Arg Val Val Gly Gly Cys Val Ala His Pro His  
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Cys Gly Gly Thr Leu Ile Ser Pro Glu Trp Val Leu Thr Ala Ala His  
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Cys Leu Glu Lys Ser Pro Arg Pro Ser Ser Tyr Lys Val Ile Leu Gly  
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Ala His Gln Glu Val Asn Leu Glu Pro His Val Gln Glu Ile Glu Val  
180 185 190

Ser Arg Leu Phe Leu Glu Pro Thr Arg Lys Asp Ile Ala Leu Leu Lys  
195 200 205

Leu Ser Ser Pro Ala Val Ile Thr Asp Lys Val Ile Pro Ala Cys Leu  
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85 90 95	
tcc aaa aca aaa aat ggc atc acc tgt caa aaa tgg agt tcc act tct Ser Lys Thr Lys Asn Gly Ile Thr Cys Gln Lys Trp Ser Ser Thr Ser	336
100 105 110	
ccc cac aga cct aga ttc tca cct gct aca cac ccc tca gag gga ctg Pro His Arg Pro Arg Phe Ser Pro Ala Thr His Pro Ser Glu Gly Leu	384
115 120 125	
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130 135 140	
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180 185 190	
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Arg Asn Pro Asp Gly Asp Val Gly Gly Pro Trp Cys Tyr Thr Thr Asn  
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Pro Arg Lys Leu Tyr Asp Tyr Cys Asp Val Pro Gln Cys Ala Ala Pro  
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Ser Phe Asp Cys Gly Lys Pro Gln Val Glu Pro Lys Lys Cys Pro Gly  
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Arg Val Val Gly Gly Cys Val Ala His Pro His Ser Trp Pro Trp Gln  
 565 570 575

Val Ser Leu Arg Thr Arg Phe Gly Met His Phe Cys Gly Gly Thr Leu  
 580 585 590

Ile Ser Pro Glu Trp Val Leu Thr Ala Ala His Cys Leu Glu Lys Ser  
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Pro Arg Pro Ser Ser Tyr Lys Val Ile Leu Gly Ala His Gln Glu Val  
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Asn Leu Glu Pro His Val Gln Glu Ile Glu Val Ser Arg Leu Phe Leu  
 625 630 635 640

Glu Pro Thr Arg Lys Asp Ile Ala Leu Leu Lys Leu Ser Ser Pro Ala  
 645 650 655

Val Ile Thr Asp Lys Val Ile Pro Ala Cys Leu Pro Ser Pro Asn Tyr  
 660 665 670

Val Val Ala Asp Arg Thr Glu Cys Phe Ile Thr Gly Trp Gly Glu Thr  
 675 680 685

Gln Gly Thr Phe Gly Ala Gly Leu Leu Lys Glu Ala Gln Leu Pro Val  
 690 695 700

Ile Glu Asn Lys Val Cys Asn Arg Tyr Glu Phe Leu Asn Gly Arg Val  
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Gln Ser Thr Glu Leu Cys Ala Gly His Leu Ala Gly Gly Thr Asp Ser  
 725 730 735

Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Phe Glu Lys Asp Lys  
 740 745 750

Tyr Ile Leu Gln Gly Val Thr Ser Trp Gly Leu Gly Cys Ala Arg Pro  
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Asn Lys Pro Gly Val Tyr Val Arg Val Ser Arg Phe Val Thr Trp Ile  
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